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FIRST NAMED INVENTOR ATTORNEY DOCKET NO. CONFIRMATION NO. APPLICATION NO. FILING DATE 01/21/2004 084335-0181 7661 10/760,320 Takao Isogai **EXAMINER** 22428 FOLEY AND LARDNER LLP MARTINELL, JAMES SUITE 500 ART UNIT PAPER NUMBER 3000 K STREET NW WASHINGTON, DC 20007 1634

DATE MAILED: 06/02/2006

Please find below and/or attached an Office communication concerning this application or proceeding.

<u> </u>				
,		Application No.	Applicant(s)	
		10/760,320	ISOGAI ET AL.	
	Office Action Summary	Examiner	Art Unit	
	-	James Martinell	1634	
Period fo	The MAILING DATE of this communication app or Reply	ears on the cover sheet with the c	orrespondence address	
WHIC - Exter after - If NO - Failui Any r	ORTENED STATUTORY PERIOD FOR REPLY CHEVER IS LONGER, FROM THE MAILING DATES as is on a firmer may be available under the provisions of 37 CFR 1.13 SIX (6) MONTHS from the mailing date of this communication. Period for reply is specified above, the maximum statutory period were to reply within the set or extended period for reply will, by statute, eply received by the Office later than three months after the mailing and patent term adjustment. See 37 CFR 1.704(b).	ATE OF THIS COMMUNICATION 36(a). In no event, however, may a reply be tim vill apply and will expire SIX (6) MONTHS from cause the application to become ABANDONE	lely filed the mailing date of this communication. (35 U.S.C. § 133).	
Status				
1)	Responsive to communication(s) filed on 28 Ma	arch 2006.		
·	• • • • • • • • • • • • • • • • • • • •	action is non-final.		
3)	Since this application is in condition for allowan	nce except for formal matters, pro	secution as to the merits is	
	closed in accordance with the practice under $\boldsymbol{\mathcal{E}}$	x parte Quayle, 1935 C.D. 11, 45	3 O.G. 213.	
Dispositi	on of Claims			
5)□ 6)⊠ 7)□	Claim(s) <u>1-14</u> is/are pending in the application.  4a) Of the above claim(s) <u>2-4 and 14</u> is/are with Claim(s) is/are allowed.  Claim(s) <u>1 and 5-13</u> is/are rejected.  Claim(s) is/are objected to.  Claim(s) are subject to restriction and/or	ndrawn from consideration.		
Applicati	on Papers			
10)🖾	The specification is objected to by the Examine The drawing(s) filed on 21 January 2004 is/are: Applicant may not request that any objection to the Carelacement drawing sheet(s) including the correction of the Oath or declaration is objected to by the Example 1.	a)⊠ accepted or b)⊡ objected drawing(s) be held in abeyance. See ion is required if the drawing(s) is obj	e 37 CFR 1.85(a). sected to. See 37 CFR 1.121(d).	
Priority u	ınder 35 U.S.C. § 119			
<ul> <li>12) Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f).</li> <li>a) All b) Some * c) None of:</li> <li>1. Certified copies of the priority documents have been received.</li> <li>2. Certified copies of the priority documents have been received in Application No</li> <li>3. Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)).</li> <li>* See the attached detailed Office action for a list of the certified copies not received.</li> </ul>				
Attachmen				
	e of References Cited (PTO-892) e of Draftsperson's Patent Drawing Review (PTO-948)	4) Interview Summary Paper No(s)/Mail Da		
3) 🔯 Inform	mation Disclosure Statement(s) (PTO-1449 or PTO/SB/08) or No(s)/Mail Date 8/04 & 8/05.		ratent Application (PTO-152)	

Art Unit: 1634

Claims 3-4 and 14 are withdrawn from further consideration pursuant to 37 CFR 1.142(b) as being drawn to a nonelected invention, there being no allowable generic or linking claim. Election was made **without** traverse in the reply filed on March 28, 2006

Amended claim 4 directed to an invention that is independent or distinct from the invention originally claimed for the following reasons: Claim 4 is drawn to the same invention as claim 3 which was Grouped in Group III in the requirement for restriction mailed November 29, 2005.

Since applicant has received an action on the merits for the originally presented invention, this invention has been constructively elected by original presentation for prosecution on the merits.

Accordingly, claim 4 is withdrawn from consideration as being directed to a non-elected invention. See 37 CFR 1.142(b) and MPEP § 821.03.

The disclosure is objected to because of the following informalities.

(a) On page 1, line 13 "arabidopsis" should be changed to "Arabidopsis" because it is a genus name.

(b) The numeral represented here by the question mark in "BRACE20353?1" on page 439, line 33 is illegible

Appropriate correction is required.

The disclosure is objected to because it contains an embedded hyperlink and/or other form of browser-executable code. Applicant is required to delete the embedded hyperlink and/or other form of browser-executable code. See MPEP § 608.01. Embedded hyperlinks and/or other forms of browser-executable code appear in at least the following locations:

- (a) page 4, lines 18-20 and 24-27,
- (b) page 9, line 28,
- (c) page 86, line 7,
- (d) page 93, line 21,
- (e) page 96, line 16,
- (f) page 98, line 20,

Art Unit: 1634

(g) page 128, lines 12, 13, and 16,

- (h) page 145, line 23,
- (i) page 151, line 27,
- (j) page 217, lines 6 and 14,
- (k) page 270, line 23, ,
- (I) page 309, lines 15 and 17, and
- (m) page 519, one 27.

The following is a quotation of the first paragraph of 35 U.S.C. 112:

The specification shall contain a written description of the invention, and of the manner and process of making and using it, in such full, clear, concise, and exact terms as to enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make and use the same and shall set forth the best mode contemplated by the inventor of carrying out his invention.

The following is a quotation of the second paragraph of 35 U.S.C. 112:

The specification shall conclude with one or more claims particularly pointing out and distinctly claiming the subject matter which the applicant regards as his invention.

Claims 1 and 5-13 are rejected under 35 U.S.C. 112, second paragraph, as being indefinite for failing to particularly point out and distinctly claim the subject matter which applicant regards as the invention. The claims are vague, indefinite, inaccurate, and misdescriptive.

- (a) Claim 1 is vague and indefinite because it claims more than was elected. It is noted that parts (b) and (c) are drawn to a non-elected invention in that the polynucleotide sequences embraced by these parts of the claim are described in terms of the amino acid sequence that is encoded. The elected invention is not described in the same terms (*i.e.*, applicants elected the nucleotide sequence SEQ ID NO: 102 and portions thereof). In addition, the claim embraces a large number of non-elected SEQ ID NOs.
- (b) The recitation of "comprising a protein-coding region" (claim 1) is vague and indefinite because the instant application does not describe the protein-coding regions of SEQ ID NO: 102.

- (c) The recitation of "which comprises the nucleotide sequence encoding a polypeptide functionally equivalent to a polypeptide encoded by [SEQ ID NO: 102]" (claim 1) is vague and indefinite because the instant application does not define the function of any polypeptide encoded by SEQ ID NO: 102 nor does it describe or define what is meant by a functional equivalent of any such putative polypeptide whose function has not been disclosed.
- (d) The recitation of "as described above" (claim 1) is vague and indefinite because "above" may describe the entire specification. Such language amounts to an omnibus-type claim.
- (e) The recitation of "comprising a nucleotide sequence encoding a partial amino acid sequence of a polypeptide encoded by the polynucleotides according to any one of (a)-(d)" is vague and indefinite. Parts (b) and (c) of the claim are drawn to a non-elected invention (the discussion in (a) above is incorporated here). The claim is also vague and indefinite because the instant application does not describe the protein-coding regions of SEQ ID NO: 102.
- (f) Claim 9 is inaccurate and misdescriptive because an oligonucleotide cannot comprise SEQ ID NO: 102 which is 3122 nucleotides in length.

Claims 1, 5-8, 12, and 13 are rejected under 35 U.S.C. 112, first paragraph, as failing to comply with the written description requirement. The claim(s) contains subject matter which was not described in the specification in such a way as to reasonably convey to one skilled in the relevant art that the inventor(s), at the time the application was filed, had possession of the claimed invention. The instant application does not describe protein-coding regions of SEQ ID NO: 102 (see the rejection under 35 U.S.C. § 112, second paragraph, items (b), (c), and (e) above). In addition, the instant application does not describe a function for any polypeptide encoded by SEQ ID NO: 102.

The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis for the rejections under this section made in this Office action:

A person shall be entitled to a patent unless -

(b) the invention was patented or described in a printed publication in this or a foreign country or in public use or on sale in this country, more than one year prior to the date of application for patent in the United States.

(e) the invention was described in (1) an application for patent, published under section 122(b), by another filed in the United States before the invention by the applicant for patent or (2) a patent granted on an application for patent by another filed in the United States before the invention by the applicant for patent, except that an international application filed under the treaty defined in section 351(a) shall have the effects for purposes of this subsection of an application filed in the United States only if the international application designated the United States and was published under Article 21(2) of such treaty in the English language.

The following is a quotation of 35 U.S.C. 103(a) which forms the basis for all obviousness rejections set forth in this Office action:

(a) A patent may not be obtained though the invention is not identically disclosed or described as set forth in section 102 of this title, if the differences between the subject matter sought to be patented and the prior art are such that the subject matter as a whole would have been obvious at the time the invention was made to a person having ordinary skill in the art to which said subject matter pertains. Patentability shall not be negatived by the manner in which the invention was made.

This application currently names joint inventors. In considering patentability of the claims under 35 U.S.C. 103(a), the examiner presumes that the subject matter of the various claims was commonly owned at the time any inventions covered therein were made absent any evidence to the contrary. Applicant is advised of the obligation under 37 CFR 1.56 to point out the inventor and invention dates of each claim that was not commonly owned at the time a later invention was made in order for the examiner to consider the applicability of 35 U.S.C. 103(c) and potential 35 U.S.C. 102(e), (f) or (g) prior art under 35 U.S.C. 103(a).

Claims 1, 5-7, and 12 are rejected under 35 U.S.C. 102(b) as being clearly anticipated by GenBank<sup>®</sup> Accession No. AC008736 (September 27, 2000). GenBank<sup>®</sup> Accession No. AC008736 has 92.7% sequence identity to SEQ ID NO: 102 (see the alignment below). Thus, GenBank<sup>®</sup> Accession No. AC008736 is embraced by the claims (*e.g.*, see Claim 1(f) and (g)). Since the DNA was sequenced, it was necessarily contained within a vector and host cell.

```
RESULT 3
AC008736/c
LOCUS
                               191925 bp
                                                           PRI 27-SEP-2000
           AC008736
                                           DNA
                                                   linear
DEFINITION Homo sapiens chromosome 19 clone CTD-2538C1, complete sequence.
ACCESSION
           AC008736
           AC008736.6 GI:10312244
VERSION
KEYWORDS
           HTG.
SOURCE
           Homo sapiens (human)
 ORGANISM Homo sapiens
           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
           Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
           Hominidae; Homo.
           1 (bases 1 to 191925)
REFERENCE
 AUTHORS
           DOE Joint Genome Institute and Stanford Human Genome Center.
  TITLE
           Direct Submission
  JOURNAL
           Unpublished
           2 (bases 1 to 191925)
REFERENCE
           DOE Joint Genome Institute.
 AUTHORS
  TITLE
           Direct Submission
  JOURNAL
           Submitted (03-AUG-1999) Production Sequencing Facility, DOE Joint
           Genome Institute, 2800 Mitchell Drive, Walnut Creek, CA 94598, USA
REFERENCE
              (bases 1 to 191925)
  AUTHORS
           DOE Joint Genome Institute and Stanford Human Genome Center.
  TITLE
           Direct Submission
           Submitted (27-SEP-2000) DOE Joint Genome Institute, 2800 Mitchell
  JOURNAL
           Drive, Walnut Creek, CA 94598, USA
COMMENT
           On Sep 27, 2000 this sequence version replaced gi:8575905.
           Draft Sequence Produced by DOE Joint Genome Institute
           www.jgi.doe.gov
           Finishing Completed at Stanford Human Genome Center
           www-shgc.stanford.edu
           Quality: Phrap Quality >=40 99.9% of Sequence;
           Estimated Total Number of Errors is 0.1.
           STS Content:
           SHGC-57769 G37408.
FEATURES
                   Location/Qualifiers
                   1. .191925
    source
                    /organism="Homo sapiens"
                    /mol type="genomic DNA"
                    /db xref="taxon:9606"
                    /chromosome="19"
                    /clone="CTD-2538C1"
ORIGIN
  Query Match
                        92.7%; Score 2892.8; DB 8; Length 191925;
                        99.9%; Pred. No. 0;
  Best Local Similarity
  Matches 2894; Conservative
                              0; Mismatches
                                                            0; Gaps
                                                                        0;
                                               2: Indels
           1 ACTAGAGGTGGGGTTAGCGCTTGGAAGCACCGACCAACGTGAGCGCAACGCGGCAGGGAC 60
Qу
             129460 ACTAGAGGTGGGGTTAGCGCTTGGAAGCACCGACCAACGTGAGCGCAACGCGGCAGGGAC 129401
Db
          61 ACCTGACCCGGGGGGCGCCCAGCCCCTCGGATTGCCAGTCACTGCTCGCTTTGGGGCACG 120
Qу
             129400 ACCTGACCCGGCGCGCCCCAGCCCCTCGGATTGCCAGTCACTGCTCGCTTTGGGGCACG 129341
Db
         121 GAGGTGCCCAGTCCTGCGGGGCACCCGACGTCCTGTCGCCGACAGGGTCCGGGAGTCAGT 180
Qу
             Db
      129340 GAGGTGCCCAGTCCTGCGGGGCACCCGACGTCCTGTCGCCGACAGGGTCCGGGAGTCAGT 129281
```

Qу	181	$\tt ATAGCTGGGTTCTAGTCCCATCACAGGCAAAAACTCCGCGGGAGCCTGGCCCGCTTTTTA$	240
Db	129280		129221
Qу	241	CCTGGGCCTCAGTTTCCCCATCCGTAAAATAGAACGGGTTGGATCTCCCGAGCGCTAACA	300
Db	129220	CCTGGGCCTCAGTTTCCCCATCCGTAAAATAGAACGGGTTGGATCTCCCGAGCGCTAACA	129161
Qу	301	$\tt TTCCAGAACTCGGATGGGCGAAGGGAGGGAGGGATGGGCCACCCAC$	360
Db	129160		129101
Qy	361	CGCGTGGAGCCCCGCCTACCACTGATCCAGGGGGTGGCAGCTCCGGCCGG	420
Db	129100	CGCGTGGAGCCCCGCCTACCACTGATCCAGGGGGTGGCAGCTCCGGCCGG	129041
Qу	421	GTGGGCGGTCCTAGGAAACCCTACCCGGCCGCCCTTGGCAGCGCCTAAGGCGGAGCGCG	480
Db	129040	GTGGGCGGGTCCTAGGAAACCCTACCCGGCCGCCCTTGGCAGCGCCTAAGGCGGAGCGCG	128981
Qу	481	CGGCTCTGCAGCCTGCTTGCCCCGGAGTTGGCACCCACGGAGGATGGGGACCGCACCCTC	540
Db	128980	CGGCTCTGCAGCCTGCTTGCCCCGGAGTTGGCACCCACGGAGGATGGGGACCGCACCCTC	128921
Qy	541	AGCTTCGCAGGGAGCCACCGTGGAGGCCAGGGCGGTGCAGAGACACGACGTGTGACTCGG	600
Db	128920	AGCTTCGCAGGGAGCCACCGTGGAGGCCAGGGCGGTGCAGAGACACGACGTGTGACTCGG	128861
Qy	601	AGTGCGCCTGGGGAGGATGGACGAGGGAGCGGGGGACCGCTAACGGGGCTCCCTCTGCGC	660
Db	128860	AGTGCGCCTGGGGAGGATGGACGAGGGAGCGGGGGACCGCTAACGGGGCTCCCTCTGCGC	128801
Qу	661	GCCCGTCCGCAGAGGCGCACGTCGAGGGTCCCGGGCGGGC	720
Db	128800	GCCCCGTCCGCAGAGGCGCACGTCGAGGGTCCCGGGCGGCTCCGTGGACGTTGGCGGTA	128741
Qу	721	GCGCCGAGCGAGTCACGGACCATGAAGAGCGTTCGTGCCGCGCGCCCAAGGCCGGGATG	780
Db	128740	GCGCCGAGCGACCATGAAGAGCGTTCGTGCCGCGCGCCCAAGGCCGGATG	128681
Qy	781	GGGGTTAGCCACATCCTGCCGCGCTGAGGGGGGGGGCCTAACGGGCGGG	840
Db	128680	GGGGTTAGCCACATCCTGCCGCGCTGAGGGGGAGGCTAACGGGCGCGGGCCGGCC	128621
Qу	841	AGCCGGAGCCCACCGCGATGGCGAGGGAGGAGTGCAAGGCGCTGCTGGACGGGCTCAACA	900
Db	128620	AGCCGGAGCCCACCGCGATGGCGAGGGAGGAGTGCAAGGCGCTGCTGGACGGGCTCAACA	128561
Qу	901	AGACGACTGCGTGCTACCACCACCTGGTGCTGACCGTCGGTGGCTCGCGGACTCGCAGA	960
Db	128560	AGACGACTGCGTGCTACCACCACCTGGTGCTGACCGTCGGTGGCTCGCCGGACTCGCAGA	128501
Qу	961	ACCTGCGGCAGAGCTGCAAAAGACGCGCCAGAAGGCGCAGGAGCTGGCGGTGTCCACCT	1020
Db	128500	ACCTGCGGCAGGAGCTGCAAAAGACGCGCCAGAAGGCGCAGGAGCTGGCGGTGTCCACCT	128441
Qу	1021	GCGCCCGGCTGACTGCTGCTGCCGCGACCGGGGCCTGGCCGCCGACGAGCGCCCGAGT	1080
Db	128440	GCGCCCGCTGACTGCTGTGCTGCCGCGACCGGGGCCTGGCCGCCGACGACGACCGAC	128381

Qγ	1081	${\tt TCGAGCGGCTCTGGGTGGCCTTCTCGGGCTGCCTGGAAGCGGACATGCGAC}$	1140
Db	128380	TCGAGCGGCTCTGGGTGGCCTTCTCGGGCTGCCTGGACCTGCTGGAAGCGGACATGCGAC	128321
Qу	1141	GCTCGCTGGAGCTGGGCGCCGCGTTCCCGCTGCACGCGCGGCGACCGCTGGTGCGCA	1200
Db	128320	GCTCGCTGGAGCTGGGCGCCGCGTTCCCGCTGCACGCGCCGCGGCGGCCGCTGGTGCGCA	128261
Qу	1201	${\tt CAGGTGTGGCGCGCCTCCTCCGGCGTGGCGCGCGCGCGCG$	1260
Db	128260	CAGGTGTGGCTGGCGCCTCCTCCGGCGTGGCGCGCGCGCG	128201
Qу	1261	GGCTCGAGGCGGAGGCGACTTCGACGTCGCGGACCTGCGGGAGCTGGAGCGCGAGGTCC	1320
Db	128200	GGCTCGAGGCGGAGCTTCGACGTCGCGGACCTGCGGAGCTGGAGCCGAGGTCC	128141
Qу	1321	TTCAGGTGGGCGAGATGATCGACAACATGGAGATGAAGGTCAACGTGCCCCGCTGGACCG	1380
Db	128140	TTCAGGTGGGCGAGATGACGACAACATGGAGATGAAGGTCAACGTGCCCCGCTGGACCG	128081
Qу	1381	TGCAAGCCCGGCAGGCGGGGCGCCGAGCTCCTGTCCACGGTCAGCGCCGGCCCCTCCT	1440
Db	128080	TGCAAGCCCGGCAGGCGCGGCGCCCTCCT	128021
Qу	1441	CGGTCGTGTCCTTGCAGGAGCGCGGGGGGGGGTTGCGACCCCAGGAAGGCCCTGGCCGCCA	1500
Db	128020	$\tt CGGTCGTGTCCTTGCAGGAGCGCGGGGGGGGGTTGCGACCCCAGGAAGGCCCTGGCCGCCA$	127961
Qу	1501	TCCTTTTCGGCGCCGTGCTGCCGGGGCTGTGGCCCTAGCCGTGTGCGTGGCGAAGCTGA	1560
Db	127960	TCCTTTTCGGCGCCGTGCTGCTGGCGGCTGTGGCCTAGCCGTGTGCGTGGCGAAGCTGA	127901
Qу	1561	GCTGACAGACACCCGACGCCGCCTGCTGCTGCCGCTCCCTGAGAAAAGACTCGG	1620
Db	127900	GCTGACAGACCCGACGCCGCCTGCTGCTGCCGCTCCCCTGAGAAAAGACTCGG	127841
Qу	1621	GATGGGTGTGGGCTGTGCAAGGGGAGTGGTCCTAAAACCCCGTGTGTGCATGG	1680
Db	127840	GATGGGTGTGGGCTGTGCAAGGGGAGTGGTCCTAAAACCCCGTGTGTGCATGG	127781
Qу	1681	GTACACGCGCGTTTCCAGTGCACATCTGCCTGGGCAGGACACGGTTTTCCTCTTGCTGGC	1740
Db	127780	GTACACGCGCGTTTCCAGTGCACATCTGCCTGGGCAGGACACGGTTTTCCTCTTGCTGGC	127721
Qу	1741	CCGGGAGAAGTTAACTTTGCGCCGGCCGTCAGGGCATTACCGCTAACGTCTGCAGGAGCT	1800
Db	127720	CCGGGAGAAGTTAACTTTGCGCCGGCCGTCAGGGCATTACCGCTAACGTCTGCAGGAGCT	127661
Qу	1801	TTATTCCCTATTAATAGAAAACCGTCACAGTGACCCTAGATCCCTCCGAGTTAATGAGTT	1860
Db	127660	TTATTCCCTATTAATAGAAAACCGTCACAGTGACCCTAGATCCCTCCGAGTTAATGAGTT	127601
Qу	1861	AACACATGTGCTGTTGGGGCGTCTTTACAGGGAGTCCGAGTTCGGTGCCCACCCCTGCCA	1920
Db	127600	AACACATGTGCTGTTGGGGCGTCTTTACAGGGAGTCCGAGTTCGGTGCCCACCCCTGCCA	127541
Qу	1921	GCGTCGCCCCTTTCTGCGTGGGACAGTTTGAAAAGGTGGGTG	1980
Db	127540	GCGTCGCCCCCTTTCTGCGTGGGACAGTTTTGAAAAGGTTGGGGTGGGGTGGAGTGAAGTTTTG	127481

Qу	1981	GAGAGGGACGCTGTTTGGTTCTATGTGGTTGGTCTGTTTCCCGGACAAGAAAATTGCAA	2040
Db	127480	GAGAGGGACGCTGTTTGGTTCTATGTGGTTGGTCTGTTTCCCGGACAAGAAAATTGCAA	127421
Qу	2041	${\tt TCAAATGTCAGCAGCTTTTATTACCTTAATCTTTCAGGGCCTAAATTTAGGAGAGTGTCC}$	2100
Db	127420	TCAAATGTCAGCAGCTTTTATTACCTTAATCTTTCAGGGCCTAAATTTAGGAGAGTGTCC	127361
Qу	2101	${\tt TGAGAGCAGTTCATACAAAGGGCTTTCTCTAAGACGCGCTACAGCCCTTCCTAGCAGAGT}$	2160
Db	127360	TGAGAGCAGTTCATACAAAGGGCTTTCTCTAAGACGCGCTACAGCCCTTCCTAGCAGAGT	127301
Qу	2161	TTATCCATTCGTCCCCAAGAGCAGCTAGAAGAGATTTGAGGTCATGACCTCCCACTGCCG	2220
Db	127300	TTATCCATTCGTCCCCAAGAGCAGCTAGAAGAGATTTGAGGTCATGACCTCCCACTGCCG	127241
Qу	2221	CTCAGGGGCTGACCCTATTTAGGAAACCAAAGAGGGTGGGT	2280
Db	127240	CTCAGGGGCTGACCCTATTTAGGAAACCAAAGAGGGTGGGT	127181
Qу	2281	TTGGATCCAGTGCGCACACTTGCCTGCGGAAAAGGGCTCTCCCCAGCCACCCGGAGATGG	2340
Db	127180	TTGGATCCAGTGCGCACACTTGCCTGCGGAAAAGGGCTCTCCCCAGCCACCCGGAGATGG	127121
Qу	2341	GGGTAAGAGGAAGAGCAGAGGCTTGGGGTAGGGCCACCTGGTGTTTAAACAGGCACTTTC	2400
Db	127120	GGGTAAGAGGAAGAGCATGGGGTAGGGCCACCTGGTGTTTAAACAGGCACTTTC	127061
Qy	2401	TCCTTCTCTGGGGCTTATTTTTGTTCAGAACTAGACCAGAGTGTTTGAACCTCCTTTGCA	2460
Db	127060	TCCTTCTCTGGGGCTTATTTTTGTTCAGAACTAGACCAGAGTGTTTGAACCTCCTTTGCA	127001
Qу	2461	GGAGGGCTGGGAATCCTCTTTAGAGCACTTAATCCTATTTATCCCCTGGAATGTGCGTGC	2520
Db	127000	GGAGGGCTGGAATCCTCTTTAGAGCACTTAATCCTATTTATCCCCTGGAATGTGCGTGC	126941
Qу	2521	TGGCCAGTAGGAGGGCTGGCTTTGGCAGCTCCCTGACCCCGCGCTGCCCGCCC	2580
Db	126940	TGGCCAGTAGGAGGCTTGGCAGCTCCCTGACCCCGCGCTGCCCGCCC	126881
Qу	2581	GGTAATGTGGCATTACTGGCCCACAGAGGTTTTGAGCCAATCAGCTCTGAGACTGGGTTA	2640
Db	126880	GGTAATGTGGCATTACTGGCCCACAGAGGTTTTGAGCCAATCAGCTCTGAGACTGGGTTA	126821
Qу		GAATGTAACAGCTTTAACTTGGGATTTAAGAAGCTTTTAAAAGGTAATAATCCTCTGAAA	
Db		GAATGTAACAGCTTTAACTTGGGATTTAAGAAGCTTTTAAAAGGTAATAATCCTCTGAAA	
Qу	2701	GAAAAATGACGTAACCACAGCGTGTACTATGAAAGCTGTTATTTTAATAAAGAACGCTGG	2760
Db		GAAAAATGACGTAACCACAGCGTGTACTATGAAAGCTGTTATTTTAATAAAGAACGCTGG	
Qу		GCCATGAACTCATACCTGCCAATGAGTCAAACATAGTATCTTTATGTAGATACTTAGATT	
Db		GCCATGAACTCATACCTGCCAATGAGTCAAACATAGTATCTTTATGTAGATACTTAGATT	
Qу		ACTAAATATATTTCATCTACTTCTGAAGTTGATAGTCTTCCCCCCCC	
Db	126640	ACTAAATATATTCATCTACTTCTGAAGTTGATAGTCTTCCCCCCCC	126581

Art Unit: 1634

Claim 8 is rejected under 35 U.S.C. 103(a) as being unpatentable over GenBank® Accession No. AC008736 (September 27, 2000) in view of applicants' admitted state of the prior art (*e.g.*, page 82, first full paragraph). GenBank® Accession No. AC008736 has 92.7% sequence identity to SEQ ID NO: 102. Applicants acknowledge the expression of nucleic acids in heterologous host cells to be old (*e.g.*, instant application at page 82, first full paragraph). It would have been obvious for one of ordinary skill in the art at the time the invention was made to express the nucleic acid of GenBank® Accession No. AC008736 in the admittedly old manner in order to produce large amounts of sequence-specific polypeptide.

Claims 1, 5-7, and 9-13 are rejected under 35 U.S.C. 102(b) as being clearly anticipated by Shoshan et al (WO 02/10449 A2 (February 7, 2002)). SEQ ID NO: 23330 of Shoshan et al is 60 nucleotides in length and matches 60 contiguous nucleotides in SEQ ID NO: 102 of the instant claims (see the alignment below). Thus, the DNA of Shoshan et al is embraced by the claims. Shoshan et al also discloses hybridization assays and antisense molecules (*e.g.*, see the abstract).

```
RESULT 14
ABN50582
ID
     ABN50582 standard; DNA; 60 BP.
XX
AC
     ABN50582;
XX
DT
     15-JUL-2002 (first entry)
XX
     Human spliced transcript detection oligonucleotide SEQ ID NO:23330.
DE
XX
     Human; mouse; rat; splice transcript; detection; RNA transcript;
KW
KW
     splice variant; transcriptome; oligonucleotide library; ss.
XX
OS
     Homo sapiens.
XX
     WO200210449-A2.
PN
XX
PD
     07-FEB-2002.
XX
     20-JUL-2001; 2001WO-IB001903.
PF
XX
     28-JUL-2000; 2000US-0221607P.
PR
     02-MAY-2001; 2001US-0287724P.
PR
XX
PΑ
     (COMP-) COMPUGEN INC.
```

```
XX
    Shoshan A, Wasserman A, Mintz E, Mintz L, Faigler S;
PΙ
XX
    WPI; 2002-257383/30.
DR
XX
PT
    New oligonucleotide libraries comprising oligonucleotides which
    selectively hybridize to mRNAs transcribed from a transcription unit of a
PT
PT
    genome, useful for detecting tissue-, pathology-, and developmental-
PΤ
    specific genes.
XX
    Example 1; SEQ ID NO 23330; 47pp; English.
PS
XX
CC
    The present invention describes oligonucleotide libraries for detecting
    messenger RNAs that populate a (sub-)transcriptome, where the (sub-
CC
    ) transcriptome comprises messenger RNAs transcribed from multiple
CC
CC
    transcription units that populate a genome. The library comprises several
    oligonucleotides, each capable of hybridising selectively to a set of
CC
CC
    messenger RNAs transcribed from a given transcription unit of the genome,
    which encodes one or more messenger RNA splice variants. The
CC
CC
    oligonucleotide libraries are useful for detecting mRNAs from a
CC
    biological sample, in expression profiling studies, in qualitatively or
    quantitatively characterising the corresponding transcriptome, and in
CC
CC
    detecting RNA transcripts and splice variants of human or animal
CC
    transcriptomes. The libraries may also be used as specialised mini
    libraries to detect transcripts of a sub-transcriptome under a particular
CC
    biological or pathological state, and so allowing the detection of tissue
CC
    - and pathology-specific genes such as those genes only expressed in
CC
    specific tissue under a specific pathological condition; to detect
CC
    developmental specific genes; and to detect RNA transcripts and splice
CC
    variants of a transcriptome of a patient suffering from a particular
CC
CC
    disorder. ABN27253 to ABN59589 represent oligonucleotide sequences from.
CC
    rats, humans and mice, which are used in the exemplification of the
    present invention. N.B. The sequence data for this patent did not form
CC
    part of the printed specification, but was obtained in electronic format
CC
    directly from WIPO at ftp.wipo.int/pub/published_pct_sequences
CC
XX
    Sequence 60 BP; 11 A; 14 C; 16 G; 19 T; 0 U; 0 Other;
SQ
  Query Match
                         1.9%; Score 60; DB 6; Length 60;
                        100.0%; Pred. No. 3.5e-17;
  Best Local Similarity
                               0; Mismatches
                                               0; Indels
                                                                Gaps
           60; Conservative
        Qу
             Db
```

Claims 1, 5-7, and 9-13 are rejected under 35 U.S.C. 102(e) as being clearly anticipated by Shoshan et al (U.S. Patent Application Publication 20030165843). SEQ ID NO: 23330 of Shoshan et al is 60 nucleotides in length and matches 60 contiguous nucleotides in SEQ ID NO: 102 of the instant claims (see the alignment below). Thus, the DNA of Shoshan et al is embraced by the claims. Shoshan et al also discloses hybridization assays and antisense molecules (*e.g.*, see the abstract).

Art Unit: 1634

```
RESULT 9
US-09-908-975-23330
; Sequence 23330, Application US/09908975
; Publication No. US20030165843A1
; GENERAL INFORMATION:
; APPLICANT: SHOSHAN, Avi
; APPLICANT: WASSERMAN, Alon
; APPLICANT: MINTZ, Eli
; APPLICANT: MINTZ, Liat
; APPLICANT: FAIGLER, Simchon
; TITLE OF INVENTION: OLIGONUCLEOTIDE LIBRARY FOR DETECTING RNA TRANSCRIPTS AND
SPLICE VARIANTS
  TITLE OF INVENTION: THAT POPULATE A TRANSCRIPTOME
  FILE REFERENCE: 36688-0005
 CURRENT APPLICATION NUMBER: US/09/908,975
 CURRENT FILING DATE: 2001-07-20
 PRIOR APPLICATION NUMBER: US 60/287,724
; PRIOR FILING DATE: 2001-05-02
; PRIOR APPLICATION NUMBER: US 60/221,607
; PRIOR FILING DATE: 2000-07-28
; NUMBER OF SEQ ID NOS: 32337
 SOFTWARE: PatentIn version 3.0
; SEQ ID NO 23330
  LENGTH: 60
   TYPE: DNA
   ORGANISM: Homo sapiens
US-09-908-975-23330
                     1.9%; Score 60; DB 3; Length 60;
 Query Match
 Best Local Similarity 100.0%; Pred. No. 3.2e-20;
                                                     0; Gaps
                                                               0;
        60; Conservative
                         0; Mismatches
                                         0; Indels
       Qу
```

Claims 1 and 5-13 are rejected under 35 U.S.C. 102(e) as being clearly anticipated by Penn et al (U.S. Patent Application Publication 20030194704). SEQ ID NO: 6988 of Penn et al is 524 nucleotides in length and matches 524 contiguous nucleotides in SEQ ID NO: 102 of the instant claims (see the alignment below). Thus, the DNA of Penn et al is embraced by the claims. Penn et al also teaches the use of arrays and nucleic acid molecular hybridization assays (*e.g.*, see paragraphs 0176-0193) and heterologous expression of nucleic acids (*e.g.*, see paragraphs 0439-0467).

```
RESULT 3
US-10-029-386-6988/c
; Sequence 6988, Application US/10029386
; Publication No. US20030194704A1
; GENERAL INFORMATION:
; APPLICANT: Penn, Sharron G.
; APPLICANT: Rank, David R.
```

```
APPLICANT: Hanzel, David K.
  TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
GENE
  TITLE OF INVENTION: EXPRESSION ANALYSIS TWO
  FILE REFERENCE: AEOMICA-X-2
  CURRENT APPLICATION NUMBER: US/10/029,386
  CURRENT FILING DATE: 2001-12-20
  NUMBER OF SEQ ID NOS: 34288
  SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
SEQ ID NO 6988
  LENGTH: 524
  TYPE: DNA
  ORGANISM: Homo sapiens
  FEATURE:
  OTHER INFORMATION: MAP TO AC010615.5
  OTHER INFORMATION: EXPRESSED IN HELA, SIGNAL = 0.56
   OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 1.2
  OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 1.3
  OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 1.1
  OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 1.5
  OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.3
  OTHER INFORMATION: EST HUMAN HIT: AW302149.1, EVALUE 0.00e+00
   OTHER INFORMATION: SWISSPROT HIT: Q06805, EVALUE 1.90e-01
   OTHER INFORMATION: NT HIT: gi4507086, EVALUE 7.80e+00
US-10-029-386-6988
                    16.8%; Score 524; DB 6; Length 524;
 Query Match
 Best Local Similarity 100.0%; Pred. No. 6.5e-265;
                                                           0;
 Matches 524; Conservative
                         0; Mismatches
                                       0;
                                         Indels
                                                     Gaps
       484 CTCTGCAGCCTGCTTGCCCCGGAGTTGGCACCCACGGAGGATGGGGACCGCACCCTCAGC 543
Qу
           524 CTCTGCAGCCTGCTTGCCCCGGAGTTGGCACCCACGGAGGATGGGGACCGCACCCTCAGC 465
Db
       544 TTCGCAGGGAGCCACCGTGGAGGCCAGGGCGGTGCAGAGACACGACGTGTGACTCGGAGT 603
Qу
           464 TTCGCAGGGAGCCACCGTGGAGGCCAGGGCGGTGCAGAGACACGACGTGTGACTCGGAGT 405
Db
       604 GCGCCTGGGGAGGATGGACGAGGGAGCGGGGACCGCTAACGGGGCTCCCTCTGCGCGCC 663
Qy
           404 GCGCCTGGGGAGGATGGACGAGGGAGCGGGGACCGCTAACGGGGCTCCCTCTGCGCGCC 345
Db
       664 CCGTCCGCAGAGGCGCACGTCGAGGGTCCCGGGCGGCTCCGTGGACGTTGGCGGTAGCG .723
Qу
           344 CCGTCCGCAGAGGCGCACGTCGAGGGTCCCGGGCGGCTCCGTGGACGTTGGCGGTAGCG 285
Db
       724 CCGAGCGAGTCACGGACCATGAAGAGCGTTCGTGCCGCGCGGCCCAAGGCCGGGATGGGG 783
Ov
           Db
       Qу
           Db
       224 GTTAGCCACATCCTGCCGCGCTGAGGGGGGGGGCCTAACGGGCCGGGCCGGGCCCAGC 165
        844 CGGAGCCCACCGCGATGGCGAGGGAGTGCAAGGCGCTGCTGGACGGGCTCAACAAGA 903
Qу
           164 CGGAGCCCACCGCGATGGCGAGGGAGGAGTGCAAGGCGCTGCTGGACGGGCTCAACAAGA 105
Db
        904 CGACTGCGTGCTACCACCTGGTGCTGACCGTCGGTGGCTCGGCGGACTCGCAGAACC 963
Qу
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Art Unit: 1634

Db 104 CGACTGCGTGCTACCACCACCTGGTGCTGACCGTCGGTGGCTCGGCGGACTCGCAGAACC 45

Qy 964 TGCGGCAGGAGCTGCAAAAGACGCGCCAGAAGGCGCAGGAGCTG 1007

Db 44 TGCGGCAGGAGCTGCAAAAGACGCGCCAGAAGGCGCAGGAGCTG 1

Any inquiry concerning this communication or earlier communications from the examiner should be directed to James Martinell whose telephone number is (571) 272-0719.

The examiner works a flexible schedule and can be reached by phone and voice mail.

Alternatively, a request for a return telephone call may be e-mailed to <a href="mailto:james.martinell@uspto.gov">james.martinell@uspto.gov</a>. Since e-mail communications may not be secure, it is suggested that information in such requests be limited to name, phone number, and the best time to return the call.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Ram Shukla, can be reached on (571) 272-0735.

Application/Control Number: 10/760,320

Art Unit: 1634

## **OFFICIAL FAX NUMBER**

The fax phone number for the organization where this application or proceeding is assigned is (571) 273-8300. Any Official Communication to the USPTO should be faxed to this number.

Any inquiry of a general nature or relating to the status of this application or proceeding should be directed to (571) 272-0547.

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James Martinell, Ph.D Primary Examiner Page 15